

Frontiers of Predictive Oncology and Computing

From Pathology to Computation The Path to Dynamic Models of Cancer

Carlos Cordon-Cardo, MD, PhD
Chair System-Wide, Department of Pathology
Mount Sinai Health System

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**Mount
Sinai**

FROM PATHOLOGY TO COMPUTATION THE PATH TO DYNAMIC MODELS OF CANCER

THEMATIC DEVELOPMENT

Sick Care → **Health Care**

Data → **Knowledge**

Treating Symptomatology → **Treating Causation**

Integration of Multidimensional Studies to Precisely Classify Disease and Optimize Treatment

Development of Comprehensive Platforms
(From Workflow and Data Flow to “Solutions-as-a-Service”)

Management of Knowledge
(From Personalized Medicine to Population Management)

Transform Practice into a Dynamic Interactive Learning Experience

A NEW PARADIGM IN PATHOLOGY

INTEGRATED APPROACH TO DISEASE MANAGEMENT

Classical Pathology  **Molecular & Systems Pathology**

Diagnosis & Staging	<p><u>Descriptive</u> analysis of clinical variables, histology, and biomarkers to categorize patients into broad disease stages; minimal to no predictive value to inform treatment selection.</p>	<p><u>Objective, quantitative</u> and multidimensional analysis of clinical variables, tissue/cellular morphometrics, and molecular signatures to define individual patient tumor phenotype and genotype, guiding treatment decisions.</p>
Prognostic Evaluation	<p>Traditional population and cohort-based classification used to deduce disease progression and likelihood of treatment response; non-specific.</p>	<p>Patient-specific characteristics and molecular tumor profiles used to predict drug sensitivity and radiation response, thus optimizing treatment efficacy and outcome.</p>
Treatment Selection	<p>Group management approach that stratifies patients into disease categories, assigning therapies on pre-determined population-based protocols instead of being patient-specific.</p>	<p>Personalized and integrated care model drives selection of evidence-based treatment protocol to optimize clinical outcome; patient-tailored treatments improve survival and quality of life.</p>

Diagnostic and prognostic approach that “groups” patients into disease categories.

Precise, predictive, and cost-effective; individualized patient management.

CENTER FOR COMPUTATIONAL AND SYSTEMS PATHOLOGY

Systems Pathology represents a novel, comprehensive approach to personalized medicine, based on the development of highly accurate predictive algorithms. Integrates clinical variables, histological/cellular features & molecular profiles through innovative technologies in the areas of image analysis, quantitative biomarker multiplexing, and deep learning.

Computational Pathology:
Mathematical characterization
of phenotype.

+

Systems Pathology:
Analytical combination of
multiple data sources.

- Board Certification in Clinical Informatics.
- Enhancing the development and funding of scientific research and clinical applications related to individualized and predictive medicine.



TRANSLATING DATA INTO KNOWLEDGE

CENTER FOR COMPUTATIONAL AND SYSTEMS PATHOLOGY

PRECISE Medical Diagnostics is a novel diagnostics platform to launch predictive/prognostic oncology-focused tests using deep learning and proprietary algorithms that combine:

- (i) Clinical features and Outcomes Data } Annotated Records
- (ii) Digital Morphology and Morphometric Statistics } Phenotype
- (iii) Molecular Pathology Signatures } Genotype
- (iv) Clinical Laboratory Parameters
- (v) Genomic Variables and Genetic Mutations

Clinical and Outcomes Data

Patient Overview Report

Patient ID: MEDKARPLE1
 Patient Date: 4/15/2017

Demographic Data

Primary Diagnosis: Breast Cancer
 Consent Date: 1/21/2017
 Date of Birth: 11/20/2017
 Sex: Female
 Race: White
 Ethnicity: Non-Hispanic
 Height (cm): 163
 Weight (kg): 65
 Body Surface Area (m²): 1.73
 Tobacco History: Current Use
 Alcohol History: Current Use

Treatment Data

Site at: MARIETTA_201481 202507 208507 201581 243502 300581 302507

Drug Administration:	Site at:	MARIETTA_201481	202507	208507	201581	243502	300581	302507
Capecitabine 250mg	mg	1	1	580	500	1	1	1
Docetaxel 75mg Sodium Phosphate	mg	1	1	1	1	1	1	1
Docetaxel 75mg NaCl	mg	50	50	50	50	50	50	50
Paclitaxel 175mg	mg	200	200	200	200	200	200	200
Zoledronic Acid 4mg	mg	1	1	1	1	1	1	1
Placebo	mg	100	100	100	100	100	100	100
Treatment	mg	200	200	200	200	200	200	200

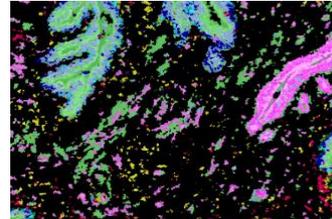
Specimen Availability

Sample Type:	Total	201481	202507	208507	201581	243502	300581	302507
ACRPTA Plasma (200uL)	17	4	3	3	3	1	1	2
Uric Acid Plasma (200uL)	12	2	2	2	2	1	1	2
ACRPTA Buffy Coat (200uL)	7	1	1	1	1	1	1	1

Morphometric Studies



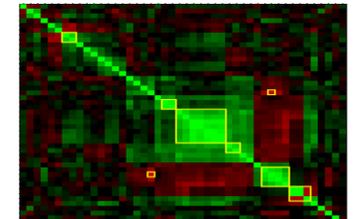
Molecular Signatures



Clinical Pathology

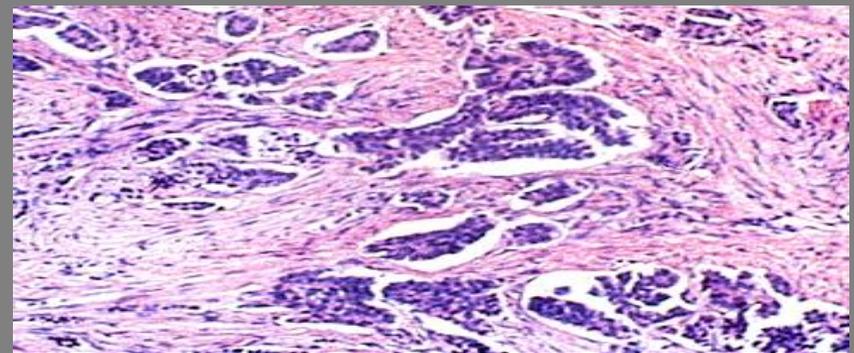
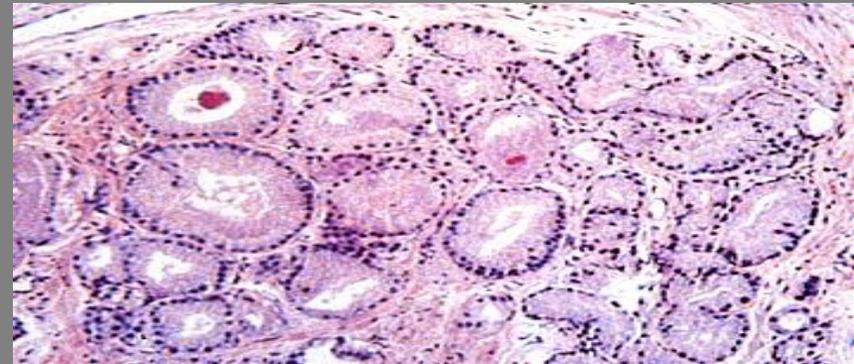
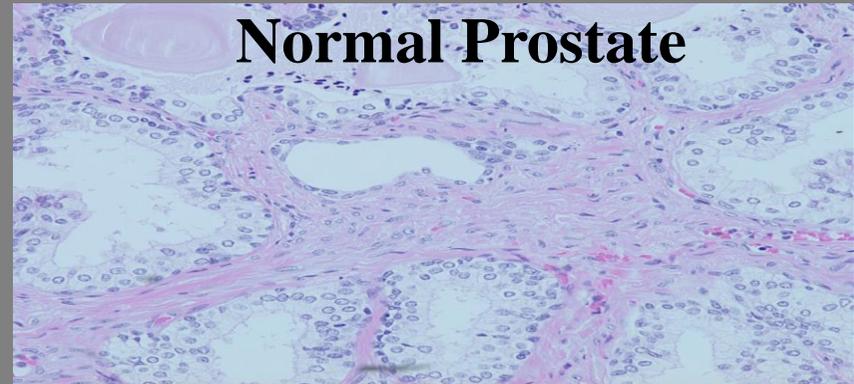
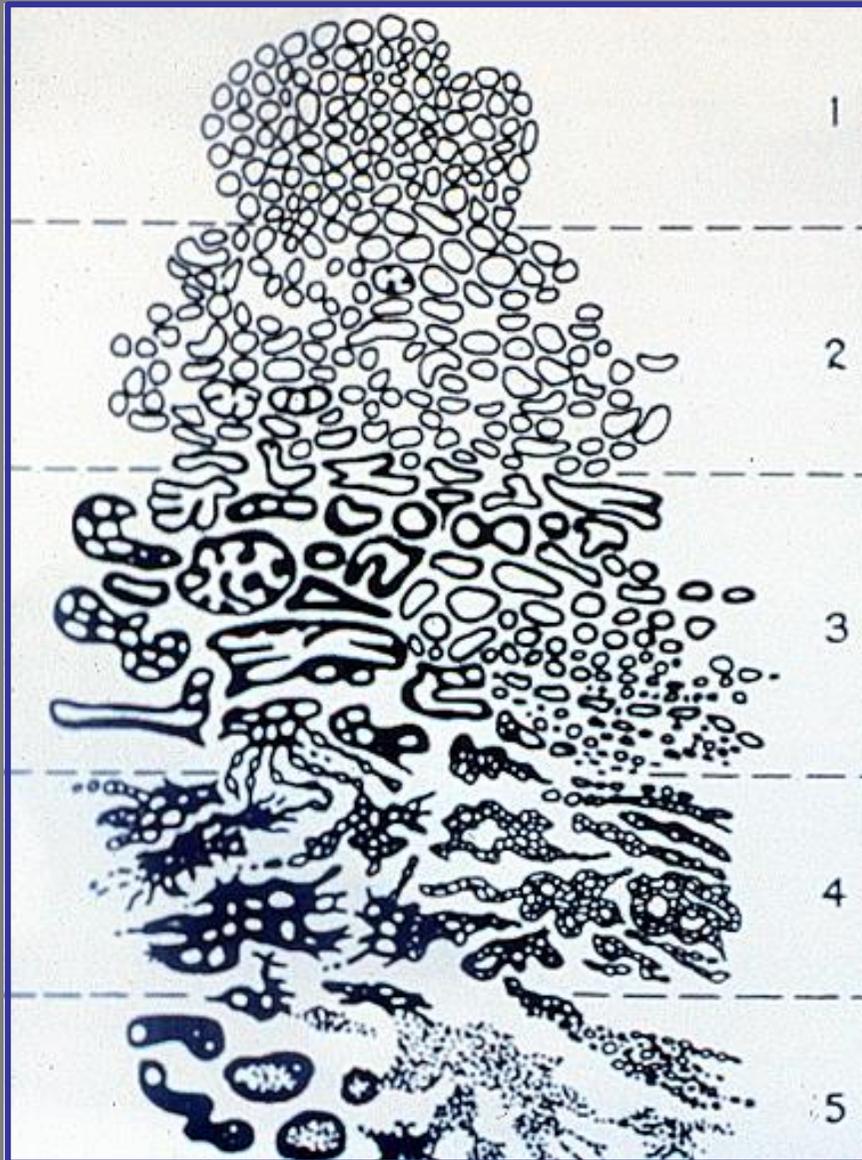


Genomic and Genetics Variables



Algorithm-Based Platform for the Accurate Diagnosis and Better Management of Cancer Patients.
 Tests with higher precision to render more effective and efficient patient care.

PROSTATE CANCER ANALYSIS: GLEASON TUMOR GRADE



PROSTATE CANCER ANALYSIS: GLEASON TUMOR GRADE

Fusion

↓

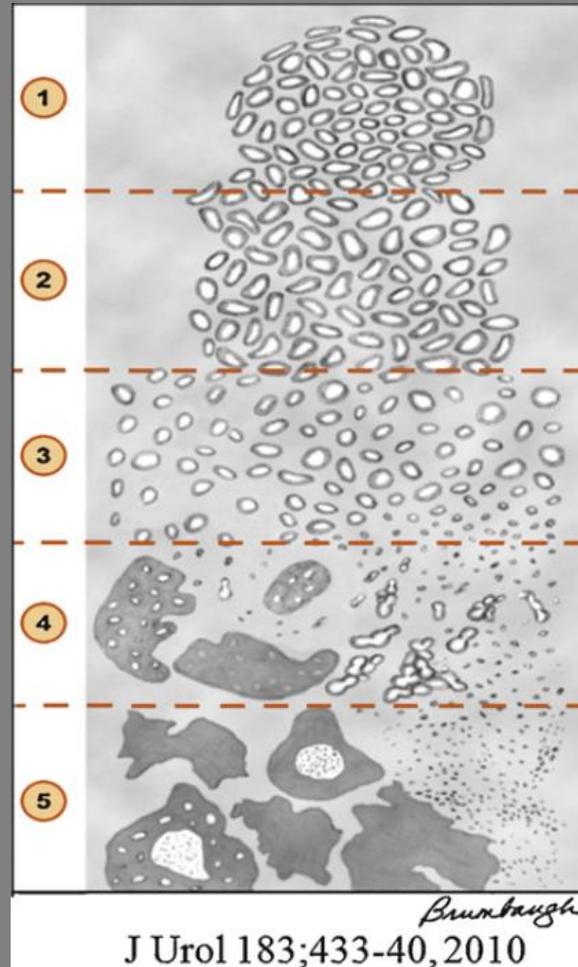
Increasing Architectural complexity

↓

Coalescent Sheets

↓

Decreasing Stromal Touch



Fragmentation

↓

Increasing Glandular & Cellular Dispersion

↓

Single Cells

↓

Decreasing Lumen Touch

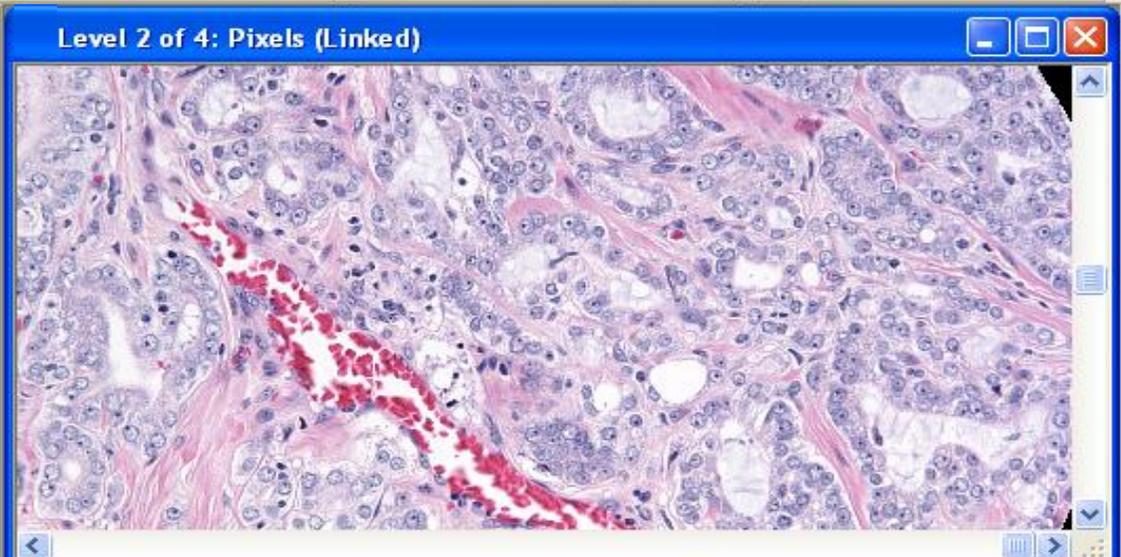
Axes of Variation and the SL Product ($S^T \cdot L^T$)



Image Object Information

Feature	Val
Process variables	
MaxBrightness	232.62
BrightnessDelta	22.38
MinBrightness	28.67
Mean_WhitespaceAr...	907.62
MinRedBlue	-16.81
GrowingStep	3.00
EndPoint	-17.68
BiggestCellSize	420.00
LargestActivePart	280.00
SmallestActivePartTo...	85.00
Red_BlueUp	1.71
Red_BlueLow	-8.68
MaxCellCompactness	1.92
Stroma/Cytoplasm thr...	3.94
New Stroma/Cytopla...	5.12
Old Stroma/Cytopla...	5.12

Features Classification

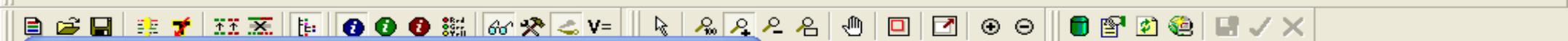


- Process
- Prostate Analysis
 - Initial Segmentation
 - Tissue / Background Analysis
 - Tissue Structure Analysis
 - Copy level 1
 - at level 2: 15 form:0.1 [c:0.9 s:0.1]
 - Classify background
 - Find dynamic thresholds
 - Find Whitespace
 - Smooth Prostatic Fluid Detection
 - Smooth Prostatic Fluid Error Correct
 - Prostatic Fluid Completed Surround
 - Lumen Next to Prostatic Fluid Detec
 - Find (RED-BLUE) threshold
 - Classify Red Blood Cells
 - Find Stroma Candidates
 - Resegment With More Weight on St
 - Find Nuclei Candidates
- Process

- Feature View
- Object features
 - Customized
 - Layer values
 - Shape
 - Generic shape features
 - Area (Px)
 - Length (Px)
 - Width (Px)
 - Length/width
 - Compactness
 - Elliptic Fit
 - Rectangular Fit
 - Border length (Px)
 - Shape index
 - Density



- Class Hierarchy
- Nucleus
 - Cytoplasm
 - Epithelial Nuclei
 - Lumen
 - Artifact
 - Stroma Nuclei
 - Cytoplasm from Stroma
 - Rejected Nuclei Candidate
 - Cytoplasm from Nuclei Candidate
 - Stroma from Nuclei Candidate
 - Red Blood Cell
 - Invading Epithelial Nuclei
- Inheritance Groups



Statistics

Settings

Image object level: Level 2

Statistic type: By Classes

Image Objects

Number Relative number (%)

Feature Operations

Sums Relative sums (%)

Means Standard deviations

Ranges

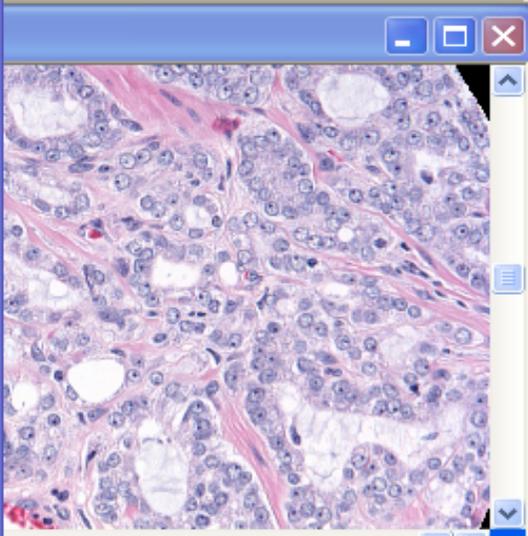
Classes

- Stroma
- Cytoplasm
- Epithelial Nuclei
- Lumen
- Artifact
- Stroma Nuclei
- Red Blood Cell

Features

- Area
- Area (excluding inner polygons)
- Area (including inner polygons)
- Asymmetry
- Average branch length
- Average Length of Branches of Order [1]
- Average length of edges (polygon)
- Avg. area represented by segments
- Border index
- Border length
- Compactness
- Compactness (foolvaon)

Save filename: Statistics



Process

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Class Hierarchy

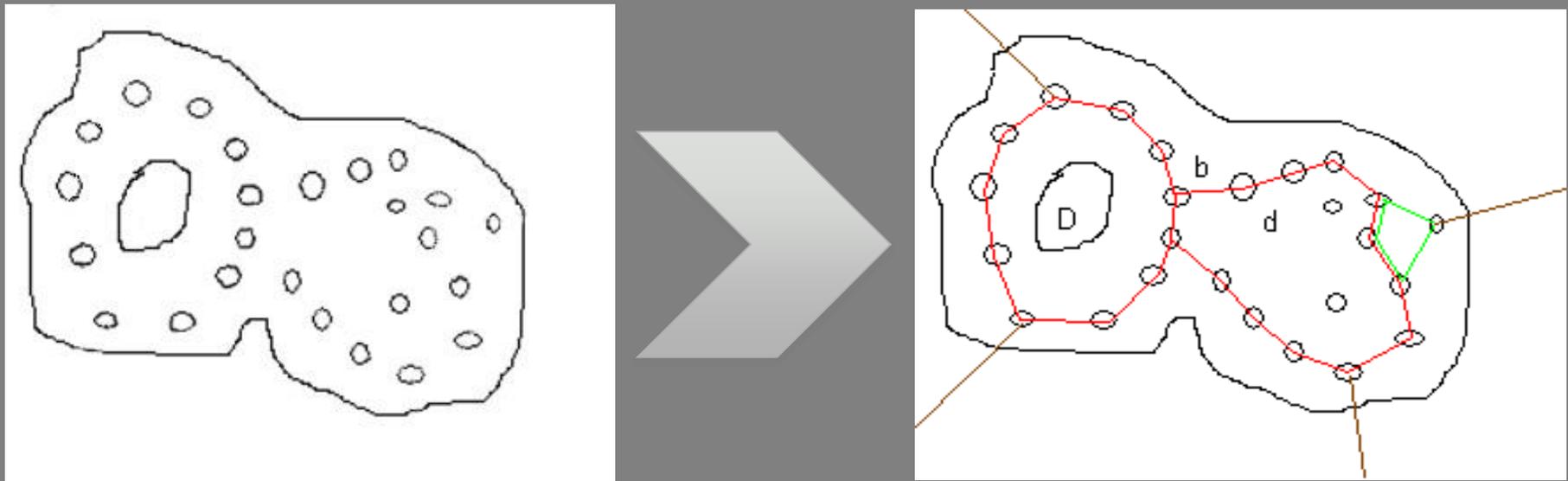
- Nucleus
- Cytoplasm

Feature Statistics

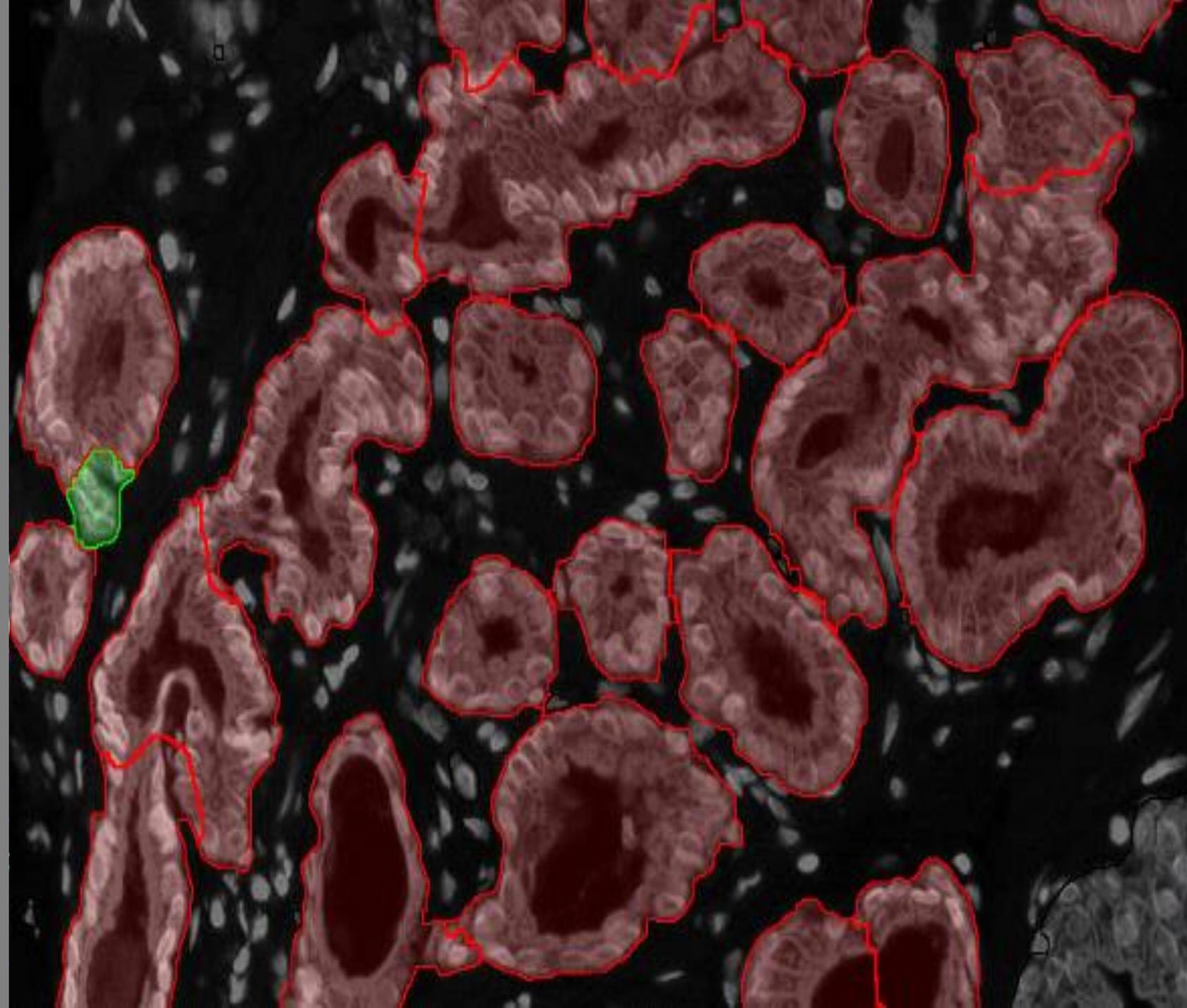
Class	Ob...	Obj...	Sum: ...	Sum r...	Mean: ...	StdDev: ...	M..	Max...	Sum: Leng...	Sum r...	Mean:...	StdDe...	M..	Max: L...
Stroma	1577	10.16	349214	19.0897	221.4420	1198.0073	1	39042	35091.4653	14.1892	22.252	31.7527	1	545.6044
Cytoplasm	9769	62.96	851292	46.5357	87.1422	106.1057	1	1957	140135.0940	56.6636	14.3449	7.8302	1	149.6356
Epithelial Nuclei	1197	7.71	341275	18.6557	285.1086	126.6422	86	1030	28348.6628	11.4628	23.683	5.9287	11	50
Lumen	91	0.59	89104	4.8709	979.1648	1242.3928	11	6417	4433.1670	1.7926	48.7161	29.6947	4	161.2188
Artifact	1546	9.96	117443	6.42	75.9657	274.9389	3	7012	22557.6623	9.1212	14.5910	16.0517	2	366.6909
Stroma Nuclei	244	1.57	49696	2.7166	203.6721	104.9983	76	574	5154.5998	2.0843	21.1254	6.7631	10	46.1819
Red Blood Cell	1093	7.04	31306	1.7113	28.6423	19.0242	2	162	11589.9252	4.6864	10.6038	4.997	2	38.6057

MORPHOMETRIC AND AUTOMATED GLEASON GRADING GLANDULAR RING STRUCTURES

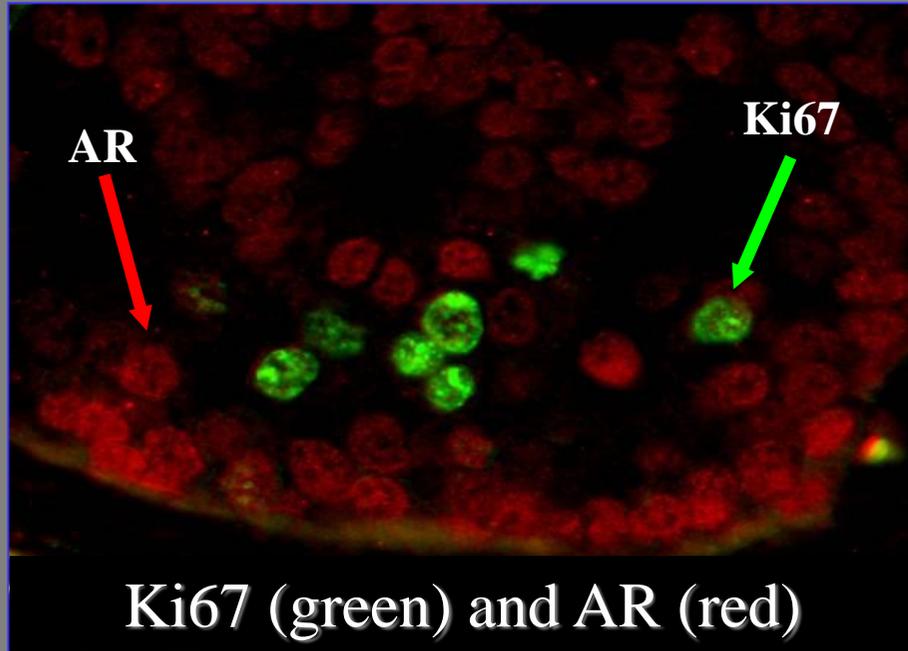
- ▶ Prostate cancer grading is based on morphological assessment of glandular differentiation (Gleason Grade).
- ▶ Gleason Grade notoriously lacks interobserver reproducibility.
- ▶ Glandular structures can be converted mathematically to “ring” structures
- ▶ Technical definition: A graph theory and “voronoi” diagram based algorithm for identifying gland rings as a basis for quantitating architectural structure, specifically degree of glandular differentiation.



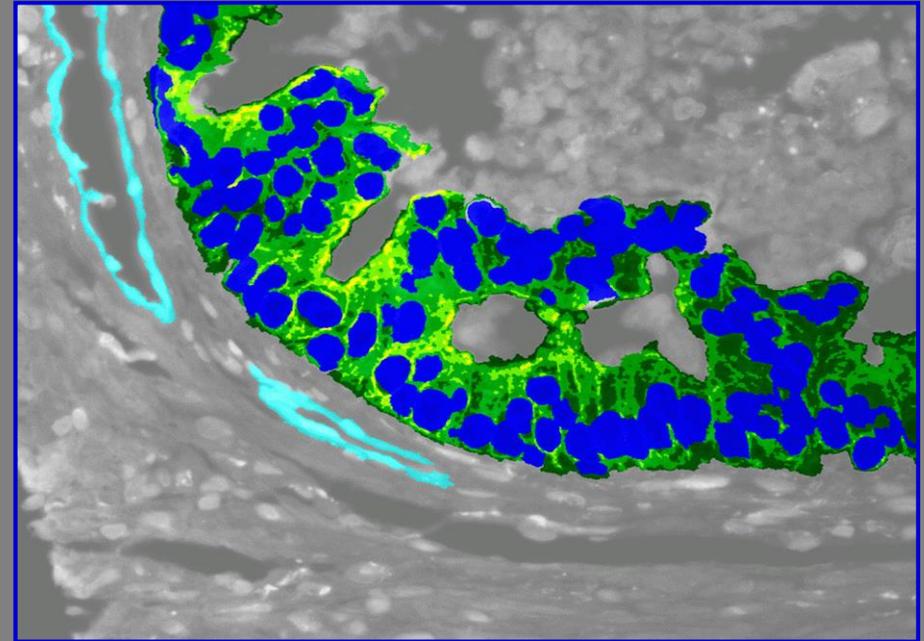
Continuous Grading (e.g., $2.4 + 2.8 = 5.2$)



PROTEIN MULTIPLEXING AT MICROANATOMICAL DETAIL



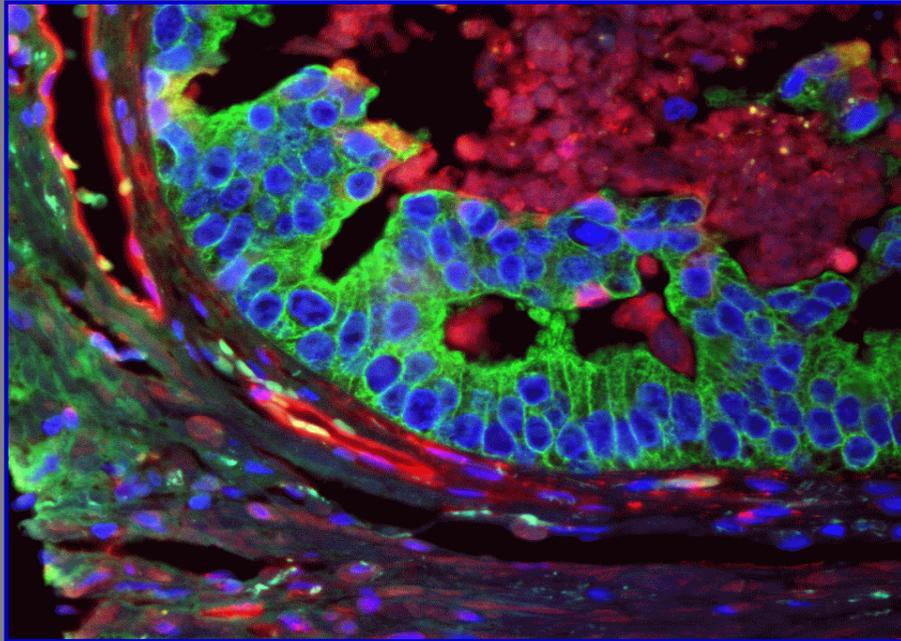
MULTIPLEXED IMAGE



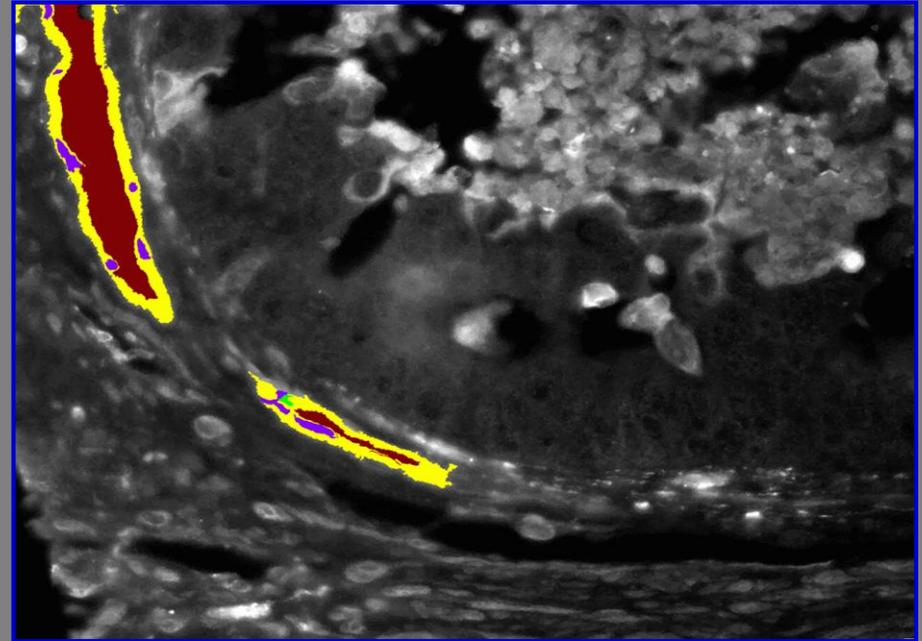
EXTRACTED FEATURE

NUMBER OF EPITHELIAL CELLS (DAPI+ CK18)
AVERAGE SIGNAL INTENSITY IN THE CYTOPLASM

PROTEIN MULTIPLEXING AT MICROANATOMICAL DETAIL



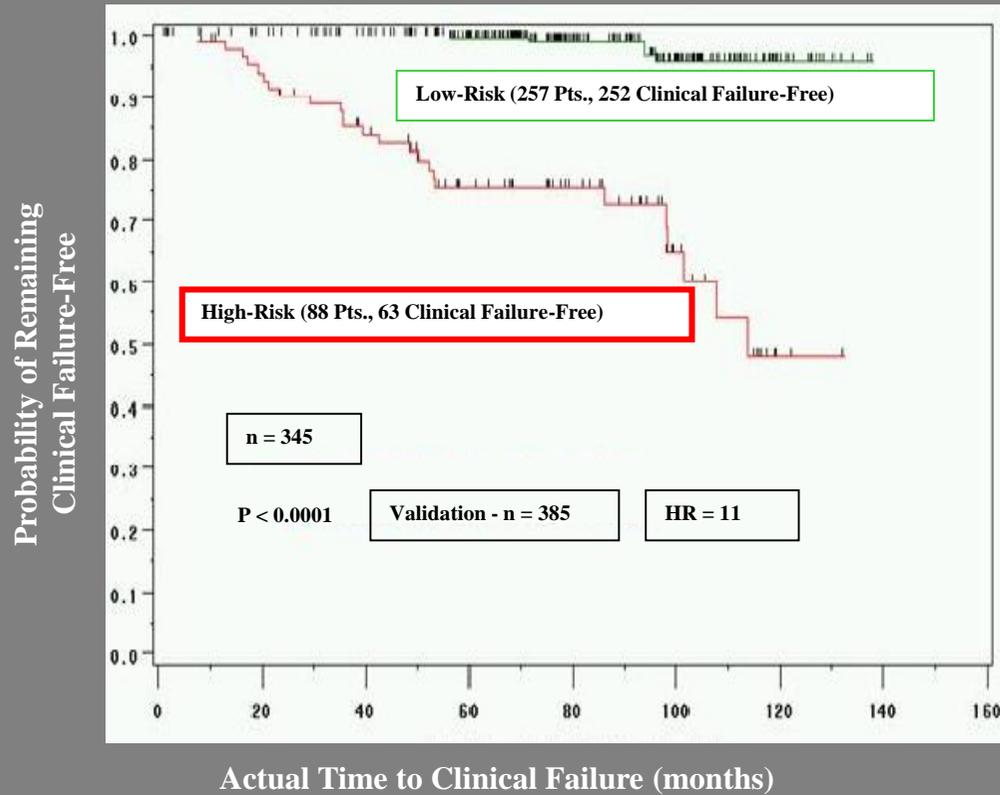
MULTIPLEXED IMAGE



EXTRACTED FEATURE

NUMBER OF LABELED VESSELS (CD34)
TOTAL VESSEL AREA, PERIMETER, LENGTH, WIDTH
MICROVESSEL AREA (MVD)

INTEGRATION OF CLINICAL AND MOLECULAR VARIABLES RENDERS PRECISE DIAGNOSIS AND PROGNOSIS



PREDICTIVE ACCURACY: 92% - SPECIFICITY: 91%; SENSITIVITY: 90%

(More than 10,000 prostate cancer cases analyzed to date)

Personalized Score Report

Precise Urology Gleason Test

Patient Information

Patient name: _____ DOB: Sex: _____
 Date of Collection: _____ MRN: _____ Specimen Type / ID: _____

Order information **Patient Clinical Details**

Order Date: _____ Specimen Receipt Date: _____ Race, Family History, PSA, _____
 Ordering Physician Name: _____ Date of last PSA, Biopsy: _____
 Ordering Physician Address: _____ Gleason grade and score: _____
 Referring Physician Name: _____

Assay Description The Precise Urology Gleason Test (PGT) analyzes a formalin-fixed paraffin-embedded (FFPE) tissue section from the original diagnostic prostate needle biopsy with a multiplex immunofluorescent assay utilizing a series of biomarkers including: AR (androgen receptor), Ki67, CK-18, CK-6 and AMACR (alpha-methylacyl-CoA racemase). Quantitative derived features reflecting histology and biology are algorithmically combined with the patient's pre-biopsy PSA and biopsy Gleason grade and score to produce a PGT score and associated probability that the patient is more or less likely to have either favorable (GS < 3+4 [ISUP ≤ 2] or unfavorable (GS ≥ 4+3 [ISUP ≥ 3] pathology in their prostatectomy specimen.

Assay Result / Interpretation:

PGT score of 12. The patient's likelihood of favorable pathology is 84% (95% CI: 76%-89%).
 Based on your patient's defined risk assessment they may be more appropriate for active surveillance.

Low Intermediate High

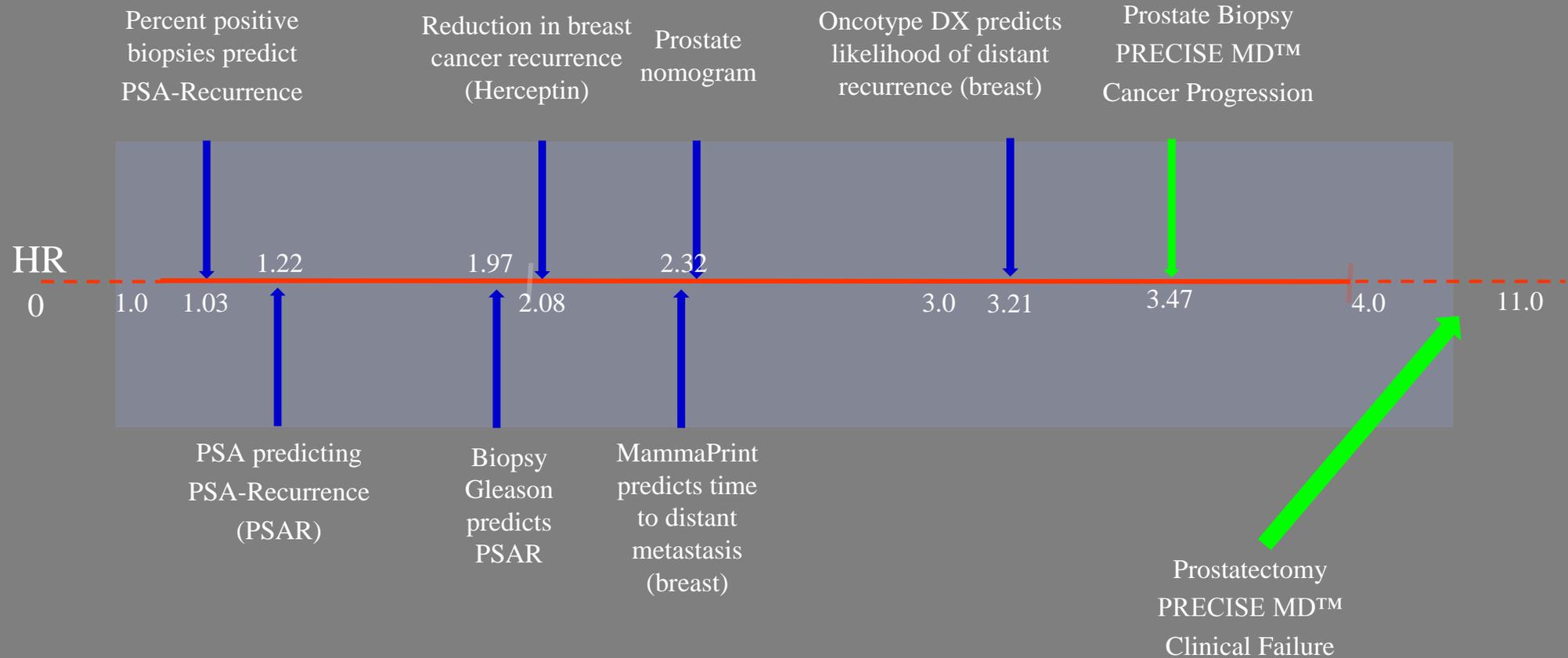
100% ← 30%

Biomarker Quantitation
AR Relative Rise Feature

Architectural Characterization
Ring Structure Feature

Physician Signature: _____ Date: _____

INTEGRATION OF CLINICAL AND MOLECULAR VARIABLES RENDERS PRECISE DIAGNOSIS AND PROGNOSIS

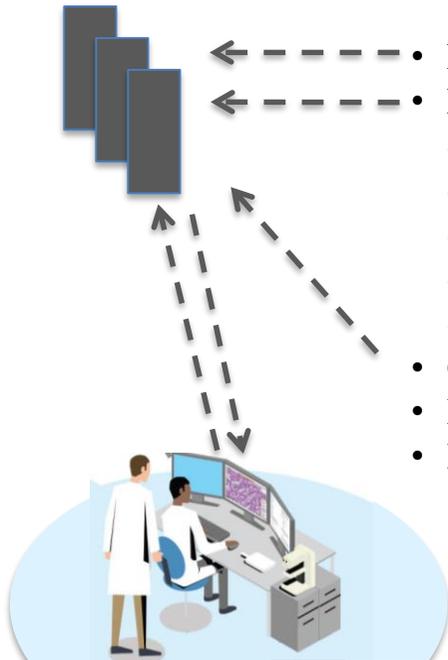


NCI-Hazard Ration (HR) (definition): A measure of how often a particular event happens in one group compared to how often it happens in another group, over time. For example, a hazard ratio of one means that there is no difference in survival between the two groups. A hazard ratio of greater than one means that survival was better in one of the groups.

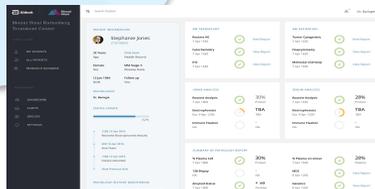
CENTER FOR COMPUTATIONAL AND SYSTEMS PATHOLOGY

DATA INPUTS

- EHR/EMR
- Laboratory LIS Result
 - Clinical Pathology
 - Anatomic Pathology
 - Molecular Pathology
 - Next Generation Sequencing
 - External Labs (e.g., media)
- Clinical Annotations/Outcomes
- Image Storage (up to 90 days)
- Image Back-up (low cost)



Digital Sign Outs



Pathology/Consoles Dashboards



Proprietary Database/Tests

CLINICAL SERVICES

CONTINUED EDUCATION AND INNOVATION

- Clinical Applications of Digital Pathology
- New Developments in Clinical Laboratory Medicine
- Clinical Informatics: A Novel Area Board Certification

GRADUATE MEDICAL EDUCATION

- Pathology & Cell/Molecular Biology Courses
 - Medical Students
 - PhD Students
- Self Assessment Modules

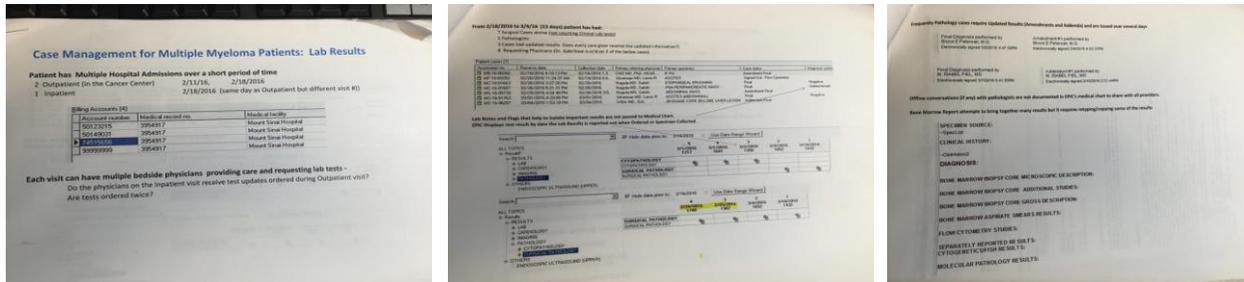
POST-GRADUATE EDUCATION

- Residency/Fellowship Training Modules
- Post-Doctoral Training Modules

EDUCATIONAL DIMENSIONS

THE NEED FOR COMPREHENSIVE PLATFORMS TEST REPORTING (MULTIPLE MYELOMA AS EXAMPLE)

FRACTIONATED and MANUAL DATA FLOW



The MM clinic manually aggregates
+20 Clinical Results per case for the
oncologist to review in 15-20 minute

CHALLENGES:

- Data residing in different systems and formats.
- Long TAT (1 to 3 wks).
- Significant manual process (risks of diagnostic errors).
- Every Hospital develops internal procedures that are not standardized and are not harmonized.

MULTIPLE MYELOMA PATHOLOGY DASHBOARD



Mount Sinai Ruttenberg
Treatment Center

QUICK LINKS

- MY PATIENTS
- ALL PATIENTS
- RESEARCH DATABASE

DASHBOARD

- DASHBOARD
- ALERTS
- GROUPS
- SETTINGS

Search Patient

Dr. Barlogie

PATIENT INFORMATION

32 Years Age [Click here](#)
Health Record

Female Sex MM Stage 3
Disease State

Follow up
Vist

PATHOLOGIST

Dr. Barlogie

STATUS UPDATE



- 1200 15 Apr 2016
Recieved Electrophoresis Results
- 600 14 Apr 2016
Avia Team
- 1300 14 Apr 2016
Patient Admitted
- [View Previous Visit](#)

PATHOLOGY HISTORY MONITORING

No major changes from pervious visit

BM TRANSPLANT

Routine HE 7 Apr 1630 [View Report](#)

Cytochemistry 7 Apr 1630 [View Report](#)

IHC 7 Apr 1630 [View Report](#)

BM ASPIRATION

Tumor Cytogetic 7 Apr 1630 [View Report](#)

Flowcytometry 7 Apr 1630 [View Report](#)

Molecular (Genseq) 7 Apr 1630 [View Report](#)

URINE ANALYSIS

Routine Analysis 7 Apr 1800 30%
Protien

Electrophoresis Eta. 9 Apr 1200 TBA
TBA

Immune Fixation NA -
NA

SERUM ANALYSIS

Routine Analysis 7 Apr 1630 28%
Protien

Electrophoresis Eta. 9 Apr 1200 TBA
TBA

Immune Fixation NA -
NA

SUMMARY OF PATHOLOGY REPORT

% Plasma Cell 7 Apr 1800 30%
Protien

138 Biopsy NA -
NA

Amyloid Status 7 Apr 1800 + ve
Positive

% Plasma on smear 7 Apr 1630 28%
Plasma

MDS 8 Apr 1200 [View Report](#)

Genetics 8 Apr 1200 [View Report](#)

THE NEED FOR COMPREHENSIVE PLATFORMS

PATHOLOGY DASHBOARDS

1. Consolidation of all Pathology Results.
2. Real-time Updates on Pathology Results.
3. Streamlined Access to stored Pathology Results.

DIRECT BENEFITS:

Reduced TAT and reporting times.
Improve Documentation/Reporting.
Translate Data into Knowledge.

INDIRECT BENEFITS:

Improves Patient Management/Outcomes.
Improves Physician Satisfaction.
Increased Accountable Care.



To assist Personalized Treatments and Improved Outcomes

INTEGRATION OF DIAGNOSTIC SERVICES: MAXIMIZING EFFECTIVENESS, IMPROVING OUTCOMES, AND MITIGATING RISK

